

SEQUENCE LISTING

<110>	Evans, Thomas Xu, Ming-Qun	
<120>	Intein-Mediated Protein Ligation Of Expressed Proteins	RECEIVE
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	09/249,543 1999-02-12	RECEIVED AUG 2 6 2003 ECH CENTER 1600/2900
<160>	26	1211/600/2900
<170>	PatentIn Ver.2.0	
<210><211><212><213>	99	
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<210><211><211><212><213>	100	T
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	4 atgat caccgtgttc tggtgatgga tggtggcctg gaatggcgtg ccgcgggtga aacgc ggcgaccgcc tggtgatgga tgatgcagct	60 100
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	5 gtttc cggcactggc aacetteegt ggcctgcgtg gegetggeeg ceaggatgtt egeta etgtttaegg tgctage	60 87
<210><211><212><212><213>	49	
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<210><211><211><212><213>	100	
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	7 recac geaggeeacg gaaggttgee agtgeeggaa actegeeage tgeateatee ragge ggtegeegeg tteeagttea eeeggggeae	60 100

<210> 8

<211><212><213>		
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	8 tccag gccaccatcc atcaccagaa cacggtgatc atgggtcaaa cgtaagcaat tcacg tgtacgcaga tcatatacgt	60 90
<210><211><211><212><213>	97	
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	9 ccaca ggtgcggaag aaacctgagg ggcatgggta gccagagccg cgaatcagtg aacgg tttgccctcc agttcagcca cagtgcg	60 97
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<223> At position 12, "Xaa" = any amino acid
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Thr Leu Glu Gly Cys Gly Glu Gln Pro Thr Gly Xaa Leu Lys
<210> 22
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                                                                        48
Gln Leu Gly Arg Ile Glu Ala Thr Asn Pro Cys Val Ser Gly Asp Thr
att gta atg aca tcc ggg ggt ccg cgg aca gtg gct gaa ctg gag ggc
                                                                        96
Ile Val Met Thr Ser Gly Gly Pro Arg Thr Val Ala Glu Leu Glu Gly
                                 25
aag ccc ttc acc gca ctt atc agg ggc tca ggg tac ccc tgc ccc tca
                                                                      144
Lys Pro Phe Thr Ala Leu Ile Arg Gly Ser Gly Tyr Pro Cys Pro Ser
        35
                            40
                                                 45
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						gaa Glu 55								192
						ttg Leu								240
						cgt Arg								288
						gct Ala								336
						gcc Ala								384
						ttc Phe 135								432
						ctc Leu							•	462
	> 1 > F > Ar	RT												
<223	Sy		size	d Fr	om M	ific Metha m.		e: Ch	emic	ally	7			

<400> 24

Gln Leu Gly Arg Ile Glu Ala Thr Asn Pro Cys Val Ser Gly Asp Thr 1 5 10 15

Ile Val Met Thr Ser Gly Gly Pro Arg Thr Val Ala Glu Leu Glu Gly 20 25 30

Lys Pro Phe Thr Ala Leu Ile Arg Gly Ser Gly Tyr Pro Cys Pro Ser 35 40 45

Gly Phe Phe Arg Thr Cys Glu Arg Asp Val Tyr Asp Leu Arg Thr Arg 50 55 60

Glu Gly His Cys Leu Arg Leu Thr His Asp His Arg Val Leu Val Met Asp Gly Gly Leu Glu Trp Arg Ala Ala Gly Glu Leu Glu Arg Gly Asp 90 85 Arg Leu Val Met Asp Asp Ala Ala Gly Glu Phe Pro Ala Leu Ala Thr 105 100 Phe Arg Gly Leu Arg Gly Ala Gly Arg Gln Asp Val Tyr Asp Ala Thr 115 120 125 Val Tyr Gly Ala Ser Ala Phe Thr Ala Asn Gly Phe Ile Val His Asn 130 135 Cys Gly Glu Gln Pro Leu Leu Thr His Glu 145 150 <210> 25 <211> 447 <212> DNA <213> Artificial Sequence <220> <221> CDS <222> (1)..(447) <223> Description of Artificial Sequence: Chemically Synthesized From Methanobacterium thermoautotrophicum. <400> 25 ctc gag gca acc aac ccc tgc gta tcc ggt gac acc att gta atg act 48 Leu Glu Ala Thr Asn Pro Cys Val Ser Gly Asp Thr Ile Val Met Thr 10 96 agt ggc ggt ccg cgc act gtg gct gaa ctg gag ggc aaa ccg ttc acc Ser Gly Gly Pro Arg Thr Val Ala Glu Leu Glu Gly Lys Pro Phe Thr 144 gca ctg att cgc ggc tct ggc tac cca tgc ccc tca ggt ttc ttc cgc Ala Leu Ile Arg Gly Ser Gly Tyr Pro Cys Pro Ser Gly Phe Phe Arg 192 acc tgt gaa cgt gac gta tat gat ctg cgt aca cgt gag ggt cat tgc Thr Cys Glu Arg Asp Val Tyr Asp Leu Arg Thr Arg Glu Gly His Cys 50 55 60 tta cgt ttg acc cat gat cac cgt gtt ctg gtg atg gat ggt ggc ctg 240

65 70 75 80	
gaa tgg cgt gcc gcg ggt gaa ctg gaa cgc ggc gac cgc ctg gtg atg Glu Trp Arg Ala Ala Gly Glu Leu Glu Arg Gly Asp Arg Leu Val Met 85 90 95	288
gat gat gca gct ggc gag ttt ccg gca ctg gca acc ttc cgt ggc ctg Asp Asp Ala Ala Gly Glu Phe Pro Ala Leu Ala Thr Phe Arg Gly Leu 100 105 110	336
cgt ggc gct ggc cgc cag gat gtt tat gac gct act gtt tac ggt gct Arg Gly Ala Gly Arg Gln Asp Val Tyr Asp Ala Thr Val Tyr Gly Ala 115 120 125	384
agc gca ttc act gct aat ggc ttc att gta cac aac tgt ggc gag cag Ser Ala Phe Thr Ala Asn Gly Phe Ile Val His Asn Cys Gly Glu Gln 130 135 140	432
cca acc ggt gaa ttc Pro Thr Gly Glu Phe 145	447
<210> 26 <211> 149 <212> PRT <213> Artificial Sequence	
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Synthesized From Methanobacterium	
Synthesized From Methanobacterium thermoautotrophicum.	
Synthesized From Methanobacterium thermoautotrophicum. <400> 26 Leu Glu Ala Thr Asn Pro Cys Val Ser Gly Asp Thr Ile Val Met Thr	
Synthesized From Methanobacterium thermoautotrophicum. <400> 26 Leu Glu Ala Thr Asn Pro Cys Val Ser Gly Asp Thr Ile Val Met Thr 1 5 10 15 Ser Gly Gly Pro Arg Thr Val Ala Glu Leu Glu Gly Lys Pro Phe Thr	
Synthesized From Methanobacterium thermoautotrophicum. <400> 26 Leu Glu Ala Thr Asn Pro Cys Val Ser Gly Asp Thr Ile Val Met Thr 1 5 10 15 Ser Gly Gly Pro Arg Thr Val Ala Glu Leu Glu Gly Lys Pro Phe Thr 20 25 30 Ala Leu Ile Arg Gly Ser Gly Tyr Pro Cys Pro Ser Gly Phe Phe Arg	
Synthesized From Methanobacterium thermoautotrophicum. <400> 26 Leu Glu Ala Thr Asn Pro Cys Val Ser Gly Asp Thr Ile Val Met Thr 1	

Asp Asp Ala Ala Gly Glu Phe Pro Ala Leu Ala Thr Phe Arg Gly Leu 100 105 110

Arg Gly Ala Gly Arg Gln Asp Val Tyr Asp Ala Thr Val Tyr Gly Ala 115 120 125

Ser Ala Phe Thr Ala Asn Gly Phe Ile Val His Asn Cys Gly Glu Gln 130 135 140

Pro Thr Gly Glu Phe 145

4 1 6